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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
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25297 7590 06/24/2009 JENKINS, WILSON, TAYLOR & HUNT, P. A. Suite 1200 UNIVERSITY TOWER 3100 TOWER BLVD., DURHAM, NC 27707				
EXAMINER				
SKOWRONEK, KARL HEINZ R				
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Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

Office Action Summary

Application No.

10/500,587

Applicant(s)

ZOU ET AL.

Examiner

KARLHEINZ R. SKOWRONEK

Art Unit

1631

Period for Reply -- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --

A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) OR THIRTY (30) DAYS, WHICHEVER IS LONGER, FROM THE MAILING DATE OF THIS COMMUNICATION.

- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.
- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.
- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

Status

- 1) ☒ Responsive to communication(s) filed on 05 May 2009.
- 2a) ☐ This action is **FINAL**. 2b) ☒ This action is non-final.
- 3) ☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11, 453 O.G. 213.

Disposition of Claims

- 4) ☒ Claim(s) 1-6, 8-13, 15-27 and 30-32 is/are pending in the application.
- 4a) Of the above claim(s) 15-27 is/are withdrawn from consideration.
- 5) ☐ Claim(s) _____ is/are allowed.
- 6) ☒ Claim(s) 1-6, 8-13 and 30-32 is/are rejected.
- 7) ☐ Claim(s) _____ is/are objected to.
- 8) ☐ Claim(s) _____ are subject to restriction and/or election requirement.

Application Papers

- 9) ☐ The specification is objected to by the Examiner.
- 10) ☐ The drawing(s) filed on _____ is/are: a) ☐ accepted or b) ☐ objected to by the Examiner.
Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).
Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).
- 11) ☐ The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.

Priority under 35 U.S.C. § 119

- 12) ☐ Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).
- a) ☐ All b) ☐ Some * c) ☐ None of:
- ☐ Certified copies of the priority documents have been received.
 - ☐ Certified copies of the priority documents have been received in Application No. _____.
 - ☐ Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).

* See the attached detailed Office action for a list of the certified copies not received.

Attachment(s)

- 1) ☐ Notice of References Cited (PTO-892)
- 2) ☐ Notice of Draftsperson's Patent Drawing Review (PTO-948)
- 3) ☒ Information Disclosure Statement(s) (PTO/SF/08)
Paper No(s)/Mail Date 05/06/09
- 4) ☐ Interview Summary (PTO-413)
Paper No(s)/Mail Date _____
- 5) ☐ Notice of Informal Patent Application
- 6) ☐ Other: _____

DETAILED ACTION

Continued Examination Under 37 CFR 1.114

A request for continued examination under 37 CFR 1.114, including the fee set forth in 37 CFR 1.17(e), was filed in this application after final rejection. Since this application is eligible for continued examination under 37 CFR 1.114, and the fee set forth in 37 CFR 1.17(e) has been timely paid, the finality of the previous Office action has been withdrawn pursuant to 37 CFR 1.114. Applicant's submission filed on 05 May 2009 has been entered.

Claim Status

Claims 1-6, 8-13, 15-27, and 30-32 are pending.

Claims 7, 14, and 28-29 are cancelled.

Claims 15-27 are withdrawn as being directed to a non-elected invention.

Claims 1-6, 8-13, and 30-32 have been examined.

Claims 1-6, 8-13, and 30-32 are rejected.

Priority

This application is the national phase application of PCT/US03/01636 filed on 17 January 2003 and claims the benefit of Provisional Application No. 60/349874 filed on 18 January 2002.

Information Disclosure Statement

The information disclosure statement (IDS) submitted on 5 May 2009 is in compliance with the provisions of 37 CFR 1.97(c). Accordingly, the information disclosure statement has been considered by the examiner.

Claim Rejections - 35 USC § 112

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

The following rejection is maintained from the previous action.

Claim 30 is rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the enablement requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to enable one skilled in the art to which it pertains, or with which it is most nearly connected, to make and/or use the invention.

In *In re Wands* (8 USPQ2d 1400 (CAFC 1988)) the CAFC considered the issue of enablement in molecular biology. The CAFC summarized eight factors to be considered in a determination of "undue experimentation." These factors include: (a) the quantity of experimentation necessary; (b) the amount of direction or guidance presented; (c) the presence or absence of working examples; (d) the nature of the invention; (e) the state of the prior art; (f) the relative skill of those in the art; (g) the predictability of the art; and (h) the breadth of the claims.

In considering the factors for the instant claims:

a) In order to use the claimed invention one of skill in the art must perform multiple hybridizations of genomic DNA to microarrays to calculate corrected hybridization signals and use the signals to calculate an expression level for a gene. For the reasons discussed below, there would be an unpredictable amount of experimentation required to practice the claimed invention.

b) The description describes determining gene expression levels from RNA samples (p.33). The specification also describes perform genomic hybridizations to identify gene probes that bind to the genomic DNA from probes that do not bind (p.23). The description does not provide detailed guidance to calculating gene expression from genomic DNA hybridization array data.

c) The description provides working examples of calculating gene expression from RNA or cDNA hybridized to arrays. The description does not provide working examples of calculating gene expression only from genomic DNA hybridized to arrays.

d) The nature of the invention, gene expression quantification, is complex.

e) The prior art does not show calculating gene expression only from genomic DNA hybridized to arrays. Kincaid et al. (Information Visualization, Vol. 4, p. 176-190, 2005) shows that genomic hybridizations to arrays measures anomalies in DNA copy number. Kincaid et al. shows genomic hybridization arrays are designed to measure the abundance of genomic DNA targets, in contrast to mRNA targets of gene expression arrays

f) The skill of those in the art of gene expression is high.

g) The predictability of calculating gene expression only from genomic DNA

hybridized to arrays is unknown in the prior art.

h) The claims are broad in that they only require genomic DNA for the determination of gene expression.

The skilled practitioner would first turn to the instant description for guidance in using the claimed invention. However, the description lacks clear evidence that genomic DNA alone can provide for the determination of gene expression. As such, the skilled practitioner would turn to the prior art for such guidance, however the prior art does not discuss the determination of gene expression levels from genomic hybridizations alone. Finally, said practitioner would turn to trial and error experimentation to determine a relationship between gene expression and genomic hybridization array data. Such amounts to undue experimentation.

Response to Arguments

Applicant's arguments filed 5 May 2009 have been fully considered but they are not persuasive. Applicant argues claim 30 has been amended to clarify that the probed are used to detect the expression level of the gene. The argument is not persuasive. Claim 30 still recites the determination of gene expression levels from the signal intensities obtained from genomic DNA as recited in claim 1 from which claim 30 depends. The rejection is maintained.

Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless –

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

The following rejection is maintained from the previous action.

Claims 1-4 are rejected under 35 U.S.C. 102(b) as being anticipated by Pietu et al. (Genome Research, Vol. 6, p. 492-503, 1996).

The claims are directed to a method of correcting oligo probe hybridization signals in which signals of multiple hybridizations are measured, a correction coefficient is calculated for each probe such that the probe's average is equal to a constant, correcting each probe signal with the correction coefficient and outputting the corrected signal.

Pietu et al. shows the analysis of hybridization signals from microarrays. Pietu et al. shows that signal intensities of multiple hybridizations are measured (p. 493, col. 2). Pietu et al. shows that each probe is divided by the average of all probes (p. 502, col.1). The correction of the signal intensity data in Pietu et al. reads on the limitations because Pietu et al. show each probe's intensity is corrected by a coefficient that is $1/(\text{average intensity})$, therefore if the average intensity is multiplied by the same coefficient the result is a constant, that is 1. This teaching reads on the limitation that each probe's signal is corrected using a coefficient such that the probe's average is equal to a constant. Pietu et al. show the probes' signal intensities are corrected and output (p. 502, col. 1). Pietu et al. shows the calculation of standard deviation and average (p.502, col. 2). Pietu et al. shows the determination of an uncertainty coefficient that is a ratio of the average to the standard deviation (p.502, col. 1).

Response to Arguments

Applicant's arguments filed 20 November 2008 have been fully considered but they are not persuasive. Applicant argues that Pietu et al. fails to teach all the elements of the claim 1, specifically a correction coefficient for each probe. The argument is not persuasive. Pietu et al. shows that to estimate the variation of measured intensities for each clone, the cv was defined as $cv = s \cdot 100/m$ where s is the standard deviation and m the mean of the values obtained in several hybridizations (p. 502, col. 1). The Calculation of CV reads on a correction coefficient as recited in the instant claims. Applicant argues that Pietu et al. do not show hybridizations using genomic DNA to correct probe hybridization. The argument is not persuasive because Pietu et al. shows that genomic or cDNA hybridizations were used to obtain signal intensity measurements that are treated as outlined on page 502. The rejection is maintained.

Claim Rejections - 35 USC § 103

The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:

(a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negated by the manner in which the invention was made.

This application currently names joint inventors. In considering patentability of the claims under 35 U.S.C. 103(a), the examiner presumes that the subject matter of the various claims was commonly owned at the time any inventions covered therein were made absent any evidence to the contrary. Applicant is advised of the obligation under 37 CFR 1.56 to point out the inventor and invention dates of each claim that was

not commonly owned at the time a later invention was made in order for the examiner to consider the applicability of 35 U.S.C. 103(c) and potential 35 U.S.C. 102(e), (f) or (g) prior art under 35 U.S.C. 103(a).

The factual inquiries set forth in *Graham v. John Deere Co.*, 383 U.S. 1, 148 USPQ 459 (1966), that are applied for establishing a background for determining obviousness under 35 U.S.C. 103(a) are summarized as follows:

1. Determining the scope and contents of the prior art.
2. Ascertaining the differences between the prior art and the claims at issue.
3. Resolving the level of ordinary skill in the pertinent art.
4. Considering objective evidence present in the application indicating obviousness or nonobviousness.

The following rejection is maintained from the previous action.

Claims 1-6 are rejected under 35 U.S.C. 103(a) as being unpatentable over Jelinsky et al. (Mol. Cell. Biol., Vol. 20, No. 21, p.8157-8167, November 2000), in view of Wohlgemuth et al. (US PG PUB 2007/0037144).

The claims are directed to a method of correcting probe hybridization signals by measuring signals from each oligo probe in multiple hybridizations; calculating a correction coefficient for the probes such that the signal average is equal to a constant; and correcting the signal for the probes using the calculated correction coefficient. In some embodiments, an average and standard deviation for the signals observed for each probe are calculated. In some embodiments, an uncertainty coefficient, called signal to noise ratio, is calculated based on the ratio of the average to standard deviation.

Jelinsky et al. teach a method of correcting oligo probe hybridization signals (p. 8157, col. 2, para 2, line 1-22). Jelinsky et al. show that 4 arrays of 6218 probes each were incubated with 10ug RNA, washed and scanned (p. 8157, col. 2, para 2, line 2-6). Jelinsky et al. show that a correction coefficient is calculated for the arrays such that the average of the intensities on the array is equal to a constant, 300 (p. 8157, col. 2, line16-18). Jelinsky et al. teach that the scaling allows the arrays to be directly compared with each other (p. 8157, col. 2, line18-19).

Jelinsky et al. do not show the calculation of individual correction coefficients for individual probes where the average signal of the individual probes is made to equal a constant.

Wohlgemuth et al. show a method of measuring DNA hybridization. Wohlgemuth et al. teach individual probes or median background subtracted signals (BGSS) can be scaled to be between 0 and 1 [0212]. Wohlgemuth et al. teach that scaling is desirable because it has the advantage of facilitating the comparison of data between different experiments [0212]. Wohlgemuth et al. teach that DNA is genomic DNA [0091]. Wohlgemuth et al. show that an average and standard deviation for the signals observed for each probe are calculated [207]. Wohlgemuth et al. show that an uncertainty coefficient, called signal to noise ratio, is calculated based on the ratio of the average to standard deviation [0207]. Wohlgemuth et al. show that probes that do not have a certain predetermined signal to noise ratio are disregarded. In Wohlgemuth et al., the signal to noise ratio is calculated as mean divided by the standard deviation. The signal to noise ratio is the inverse of the coefficient of variation, standard deviation

divided by the mean. Wohlgemuth et al. teach that if the signal to noise ratio is less than 3 which is equivalent to having a coefficient of variation that is greater than 0.33 the data is flagged but used, reading on a predetermined value that is approximately 1.0 [0728]. As the signal to noise ratio decreases, it approaches 1. When the signal to noise ratio is equal to 1, the ratio indicates that signal cannot be distinguished from the noise, indicative of data of poor quality. Similarly, for the inverse of the signal to noise ratio, coefficient of variation (CV), as the CV increases to approach 1, the quality of the data becomes poorer and less reliable. Wohlgemuth et al. teach that if a replicate feature is of poor quality it can be disregarded and the remaining features used to represent the gene [0200].

It would have been obvious to one of ordinary skill in the art at the time of invention to modify the method of correcting oligo probe hybridization signals of Jelinsky et al. with the individual probe scale factors of Wohlgemuth et al. because Wohlgemuth et al. shows that scaling provides the advantage of facilitating the comparison of data between different experiments. As the signal to noise ratio decreases, it approaches 1. When the signal to noise ratio is equal to 1, the ratio indicates that signal cannot be distinguished from the noise, indicative of data of poor quality. Similarly, for the inverse of the signal to noise ratio, coefficient of variation (CV), as the CV increases to approach 1, the quality of the data becomes poorer and less reliable.

Response to Arguments

Applicant's arguments filed 05 May 2009 have been fully considered but they are not persuasive. In response to applicant's arguments against the references individually,

one cannot show nonobviousness by attacking references individually where the rejections are based on combinations of references. See *In re Keller*, 642 F.2d 413, 208 USPQ 871 (CCPA 1981); *In re Merck & Co.*, 800 F.2d 1091, 231 USPQ 375 (Fed. Cir. 1986). Applicant argues that the recitation of genomic DNA in [0091] of Wohlgemuth et al. is taken out of context. The argument is not persuasive. In Paragraph [0091], Wohlgemuth et al. shows DNA molecules may be genomic DNA and that one can identify sequences of interest for analyzing gene expression using the procedures disclosed. This is a direct suggestion by Wohlgemuth et al. that genomic DNA may be used in the processes disclosed in [0207-0217]. Applicant argues that the data scaling in [0212] is not the same as correcting probe hybridization signals because it based on the mean of the entire dataset. This is not persuasive. Wohlgemuth et al. points out in [0214] that each probe or individual expression level is corrected using a scaling factor. The rejection is maintained.

The following rejection is maintained from the previous action.

Claim 8-13 and 31-32 are rejected under 35 U.S.C. 103(a) as being unpatentable over Jelinsky et al. (Mol. Cell. Biol., Vol. 20, No. 21, p.8157-8167, November 2000), in view of Wohlgemuth et al. (US PG PUB 2007/0037144) as applied to claims 1-5 above, and further in view of Pinkel et al. (US Pat 5,830,645).

The claims are drawn to determining a dynamic range for DNA binding.

Jelinsky et al. (Mol. Cell. Biol., Vol. 20, No. 21, p.8157-8167, November 2000), in view of Wohlgemuth et al. (US PGPUB 2007/0037144) as applied to claims 1-5 above do not teach determining a dynamic range for DNA binding.

Pinkel et al. show a method of comparative genomic hybridization. Pinkel et al. show that the method provides increased sensitivity, more precise localization of chromosomal abnormalities and which can detect differences in levels of gene expression are particularly desirable for the diagnosis of disease (col. 2, line 23-26). Pinkel et al. show that serial dilutions of pairs of fluorochrome in known relative proportions can also be analyzed to determine the accuracy with which fluorescence ratio measurements reflect actual fluorochrome ratios over the dynamic range permitted by the detectors and membrane fluorescence (col. 8, line 44-49).

It would have been obvious to one of ordinary skill in the art at the time of invention to modify the method of correcting oligo probe hybridization signals of Jelinsky et al., in view of Wohlgemuth et al. as applied to claims 1-5 above and in further view of the method comparative genomic hybridization by Pinkel et al. because Pinkel et al. show that the method provides increased sensitivity, more precise localization of chromosomal abnormalities and which can detect differences in levels of gene expression are particularly desirable for the diagnosis of disease.

Response to Arguments

Applicant's arguments filed 5 May 2009 have been fully considered but they are not persuasive. Applicant argues that Pinkel et al. does not cure the deficiencies of

Jelinsky et al. in view of Wohlgemuth et al. The argument is not persuasive for the reasons given above. The rejection is maintained.

Conclusion

Any inquiry concerning this communication or earlier communications from the examiner should be directed to KARLHEINZ R. SKOWRONEK whose telephone number is (571)272-9047. The examiner can normally be reached on 8:00am-5:00pm Monday-Friday.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Marjorie Moran can be reached on (571) 272-0720. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see <http://pair-direct.uspto.gov>. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free). If you would like assistance from a USPTO Customer Service Representative or access to the automated information system, call 800-786-9199 (IN USA OR CANADA) or 571-272-1000.

/KARLHEINZ R SKOWRONEK/
Examiner, Art Unit 1631

24 June 2009